



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/511527

Source:

PCT

Date Processed by STIC:

10-26-04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
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Revised 05/17/04



PCT

RAW SEQUENCE LISTING

DATE: 10/26/2004

PATENT APPLICATION: US/10/511,527

TIME: 17:53:54

Input Set : A:\231258.st25.txt

Output Set: N:\CRF4\10262004\J511527.raw

4 <110> APPLICANT: Brandt Dr., Burkhard H.
 6 <120> TITLE OF INVENTION: Method for characterizing primary tumours
 W--> 7 <130> FILE REFERENCE: PCT/EP03/04037
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/511,527
 C--> 9 <141> CURRENT FILING DATE: 2004-10-15
 9 <160> NUMBER OF SEQ ID NOS: 42
 11 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply
 Corrected Diskette Needed

(PS: 1-3) ←

ERRORED SEQUENCES

178 <210> SEQ ID NO: 16
 179 <211> LENGTH: 20
 180 <212> TYPE: DNA
 E--> 181 <213> Artificial Sequence
 183 <220> FEATURE:
 184 <223> OTHER INFORMATION: Description of the artificial Sequence: Oligonucleotide
 E--> 186 <213> ORGANISM: ignore
 186 <400> SEQUENCE: 18
 187 ctgggagtca agtgttttgg 20
 200 <210> SEQ ID NO: 18
 201 <211> LENGTH: 18
 202 <212> TYPE: DNA
 E--> 203 <213> Artificial Sequence
 205 <220> FEATURE:
 206 <223> OTHER INFORMATION: Description of the artificial Sequence: Oligonucleotide
 E--> 208 <213> ORGANISM: ignore
 208 <400> SEQUENCE: 18
 209 aacggtcatc tgtgttcg 18
 222 <210> SEQ ID NO: 20
 E--> 223 <211> 20
 224 <212> TYPE: DNA
 225 <213> ORGANISM: Artificial Sequence
 227 <220> FEATURE:
 228 <223> OTHER INFORMATION: Description of the artificial Sequence: Oligonucleotide
 E--> 230 <211> LENGTH:
 230 <400> SEQUENCE: 20
 OK E--> 231 ttgctgtag aggatgcaat 20

Replace lowercase letter "L" with numeral 1.

Same error

Same error

Same error

<210> 1
<211> 20
<212> DNA

<213> Artificial Sequence

PLS explain source of genetic material.

<220>

<223> Description of the artificial sequence

Invalid response

<400> 1

gcaggacatg .agatgactga

20

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/511,527

DATE: 10/26/2004

TIME: 17:53:55

Input Set : A:\231258.st25.txt

Output Set: N:\CRF4\10262004\J511527.raw

L:7 M:283 W: Missing Blank Line separator, <130> field identifier
L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:181 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER
L:186 M:282 E: Numeric Field Identifier Missing, <213> is required
L:203 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER
L:208 M:282 E: Numeric Field Identifier Missing, <213> is required.
L:223 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER
L:230 M:282 E: Numeric Field Identifier Missing, <211> is required.
L:231 M:252 E: No. of Seq. differs, <211> LENGTH:Input:0 Found:20 SEQ:20